

Db 20 AlaGluSerTrpAspAsnValGlyLeu 29

RESULT 2
US-09-328-352-7079
; Sequence 7079, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352
; CURRENT FILING DATE: 1999-06-04
; NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO 7079
; LENGTH: 289
; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-09-328-352-7079

Alignment Scores:
Pred. No.: 23.6 Length: 289
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.61% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-09-328-352-7079 (1-289)

OY 384 GTCAGTCATGTCAGGAGAGT 361
Db 11 ValArgSerLeuValArgLysSer 18

RESULT 3
US-08-709-838-2
; Sequence 2, Application US/08709838
; Patent No. 6140064
; GENERAL INFORMATION:
; APPLICANT: Loetscher, Marcel
; APPLICANT: Moser, Bernhard
; TITLE OF INVENTION: IP-10/MIG RECEPTOR DESIGNATED CXCR3,
; TITLE OF INVENTION: NUCLEIC ACIDS, AND METHODS OF USES THEREFOR
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
; STREET: Two Militia Drive
; CITY: Lexington
; STATE: MA
; COUNTRY: USA
; ZIP: 02173
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/709,838
; FILING DATE:
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: Brook Esq., David E.
; REGISTRATION NUMBER: 22,592
; REFERENCE/DOCKET NUMBER: TK196-01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 861-6240
; TELEFAX: (617) 861-9540
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 368 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-709-838-2

Alignment Scores:
Pred. No.: 22.8 Length: 368
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.58% Indels: 0
DB: 3 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-709-838-2 (1-368)

OY 744 TCACCTCTTTCTCTAGACTG 767
Db 61 SerLeuPheLeuLeuGlyLeu 68

RESULT 4
US-08-829-839-2
; Sequence 2, Application US/08829839
; Patent No. 6184358
; GENERAL INFORMATION:
; APPLICANT: Loetscher, Marcel
; APPLICANT: Moser, Bernhard
; APPLICANT: Qin, Shixin
; APPLICANT: Mackay, Charles R.
; TITLE OF INVENTION: IP-10/MIG RECEPTOR DESIGNATED CXCR3,
; TITLE OF INVENTION: ANTIBODIES, NUCLEIC ACIDS, AND METHODS OF USE THEREFOR
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
; STREET: Two Militia Drive
; CITY: Lexington
; STATE: MA
; COUNTRY: USA
; ZIP: 02173
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/829,839
; FILING DATE: 31-MAR-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/709,838
; FILING DATE: 10-SEP-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Brook Esq., David E.
; REGISTRATION NUMBER: 22,592
; REFERENCE/DOCKET NUMBER: TK196-01A
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 861-6240
; TELEFAX: (617) 861-9540
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 368 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-829-839-2

Alignment Scores:
Pred. No.: 22.8 Length: 368
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.58% Indels: 0
DB: 3 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-829-839-2 (1-368)

OY 744 TCACCTCTTTCTCTAGACTG 767
Db 61 SerLeuPheLeuLeuGlyLeu 68

```
RESULT 5
US-09-170-496D-20
; Sequence 20, Application US/09170496D
; Patent No. 6555339
; GENERAL INFORMATION:
; APPLICANT: Behan, Dominic P.
; APPLICANT: Chalmers, Derek T.
; APPLICANT: Liaw, Chen W.
; TITLE OF INVENTION: No. 6555339-Endogenous, Constitutively Activated Human G Protein-
; TITLE OF INVENTION: Receptors
; FILE REFERENCE: AREN-0040
; CURRENT APPLICATION NUMBER: US/09/170,496D
; CURRENT FILING DATE: 1998-10-13
; NUMBER OF SEQ ID NOS: 294
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 368
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-170-496D-20

Alignment Scores:
Pred. No.: 22.8 Length: 368
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.58% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-09-170-496D-20 (1-368)
Oy 744 TCACCTCTTTTCTGCTAGAGCTG 767
Db 61 SerleuLeupheLeuLeuGlyLeu 68

RESULT 6
US-09-170-496D-174
; Sequence 174, Application US/09170496D
; Patent No. 6555339
; GENERAL INFORMATION:
; APPLICANT: Behan, Dominic P.
; APPLICANT: Chalmers, Derek T.
; APPLICANT: Liaw, Chen W.
; TITLE OF INVENTION: No. 6555339-Endogenous, Constitutively Activated Human G Protein-
; TITLE OF INVENTION: Receptors
; FILE REFERENCE: AREN-0040
; CURRENT APPLICATION NUMBER: US/09/170,496D
; CURRENT FILING DATE: 1998-10-13
; NUMBER OF SEQ ID NOS: 294
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 174
; LENGTH: 368
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-170-496D-174

Alignment Scores:
Pred. No.: 22.8 Length: 368
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.58% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-09-170-496D-174 (1-368)
Oy 744 TCACCTCTTTTCTGCTAGAGCTG 767
Db 61 SerleuLeupheLeuLeuGlyLeu 68

RESULT 7
US-09-252-991A-24098
; Sequence 24098, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 24098
; LENGTH: 509
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-24098

Alignment Scores:
Pred. No.: 21.9 Length: 509
Score: 8.00 Matches: 8
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.58% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-09-252-991A-24098 (1-509)
Oy 744 TCACCTCTTTTCTGCTAGAGCTG 767
Db 188 SerleuLeupheLeuLeuGlyLeu 195

RESULT 8
US-08-340-283-19
; Sequence 19, Application US/08340283
; Patent No. 5861318
; GENERAL INFORMATION:
; APPLICANT: Elhammer, Ake P.
; TITLE OF INVENTION: A SCINTILLATION PROXIMITY ASSAY FOR
; TITLE OF INVENTION: N-ACETYLGLACTOSAMINYLTRANSFERASE ACTIVITY
; NUMBER OF SEQUENCES: 205
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pharmacia and Upjohn, Inc., Intellect. Prop. Law
; ADDRESSSEE: (1920-32-1)
; STREET: 301 Henrietta Street
; CITY: Kalamazoo
; STATE: Michigan
; COUNTRY: U.S.A.
; ZIP: 49001
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/340,283
; FILING DATE:
; CLASSIFICATION: 436
; ATTORNEY/AGENT INFORMATION:
; NAME: Woolton, Thomas A.
; REGISTRATION NUMBER: 35,004
; REFERENCE/DOCKET NUMBER: 4828
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (616) 385-7914
; TELEFAX: (616) 385-6897
; TELEX: 224401
; INFORMATION FOR SEQ ID NO: 19:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 9 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
```

TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
US-08-340-283-19

Alignment Scores:
Pred. No.: 9.27e+06 Length: 9
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 2 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-340-283-19 (1-9)

OY 407 GCACCTCCTGCATCCTCCT 387
Db 1 AlapropioProSerLeuPro 7

RESULT 9
US-10-053-485-46
Sequence 46, Application US/10053485
Patent No. 6576896
GENERAL INFORMATION:
APPLICANT: Figeys, Daniel
APPLICANT: Aebersold, Ruedi
TITLE OF INVENTION: ELECTROOSMOTIC FLUIDIC DEVICE AND RELATED METHODS
FILE REFERENCE: UMOCT118617
CURRENT APPLICATION NUMBER: US/10/053,485
CURRENT FILING DATE: 2002-05-28
PRIOR APPLICATION NUMBER: US 09/209,880
PRIOR FILING DATE: 1998-12-11
PRIOR APPLICATION NUMBER: US 60/069,398
PRIOR FILING DATE: 1997-12-12
NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn version 3.0
SEQ ID NO 46
LENGTH: 12
TYPE: PPT
ORGANISM: Saccharomyces cerevisiae
US-10-053-485-46

Alignment Scores:
Pred. No.: 348 Length: 12
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.38% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-10-053-485-46 (1-12)

OY 1193 CGAGCTTCTTCTGCACCTT 1213
Db 1 AtgGlypheLeuSerAspLeu 7

RESULT 10
US-08-908-371B-9
Sequence 9, Application US/08908371B
Patent No. 6331610
GENERAL INFORMATION:
APPLICANT: Bourinbalar, Aldar S.
TITLE OF INVENTION: A Method for Preventing and Treating
TITLE OF INVENTION: AIDS and HIV Infection Using Select Peptides From the
TITLE OF INVENTION: Beta Subunit of Human Chorionic Gonadotropin
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Metation, Inc.
STREET: 367 Bay Shore Road
CITY: Deer Park
STATE: New York

COUNTRY: United States of America
ZIP: 11729
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 in. diskette (1.44megabytes)
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: WORD 6.0 ASCII TEXT CONVERSION ONLY
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/908,371B
FILING DATE: 07-AUG-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/044,937
FILING DATE: 25-APR-1997
ATTORNEY/AGENT INFORMATION:
NAME: COLEMAN, HENRY D.
REGISTRATION NUMBER: 32,559
REFERENCE/DOCKET NUMBER: M31-013
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 679-0090
TELEFAX: (212) 679-9121
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acid units
type: Amino Acid
STRANDEDNESS: Single Stranded
TOPOLOGY: N/A
MOLECULE TYPE: Protein Subunit
DESCRIPTION: 13 amino acids corresponding to amino
DESCRIPTION: acids 120-132 of the beta subunit of human chorionic
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE: Chemical Synthesis
IMMEDIATE SOURCE: N/A
POSITION IN GENOME: N/A
FEATURE:
NAME/KEY: Oligopeptide corresponding to Amino Acids 120-132
NAME/KEY: of Beta Subunit of Human Chorionic Gonadotropin
LOCATION: Beta subunit of Human Chorionic Gonadotropin
IDENTIFICATION METHOD: Chemical synthesis
PUBLICATION INFORMATION:
AUTHORS: STEVENS, Vernon C.
TITLE: ANTIGENIC MODIFICATION OF AMINO ACIDS
JOURNAL: UNITED STATES PATENT NUMBER 4,302,386
VOLUME: N/A
PAGES: N/A
DATE: No. 6331610ember 24, 1981
US-08-908-371B-9

Alignment Scores:
Pred. No.: 345 Length: 13
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-908-371B-9 (1-13)

OY 407 GCACCTCCTGCATCCTCCT 387
Db 4 AlapropioProSerLeuPro 10

RESULT 11
US-08-569-147-57
Sequence 57, Application US/08569147
Patent No. 6180377
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: HUMANISED ANTIBODIES
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:

ADDRESSEE: Woodcock Washburn Kurtz MacKiewicz &
ADDRESSEE: No. 6180377rls, LLP
STREET: One Liberty Place - 46th Floor
City: Philadelphia
STATE: PA
COUNTRY: U.S.A.
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/569,147
FILING DATE: 25-March-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yalko
REGISTRATION NUMBER: 35,719
REFERENCE/DOCKET NUMBER: CARP-0047
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 57:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-569-147-57

Alignment Scores:
Pred. No.: 341 Length: 14
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.38% Indels: 0
DB: 3 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-569-147-57 (1-14)

OY 145 CCTGGAAGAGCCGTGAAGTGG 165
DB 6 Proglylsglyleuylstrp 12

RESULT 12
5451527-3
Patent No. 5451527
APPLICANT: SARIN, VIRENDER K.;BODNER, JOHN B.
TITLE OF INVENTION: HCG PEPTIDES FOR USE IN ANTIBODY
PURIFICATION PROCEDURES
NUMBER OF SEQUENCES: 14
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/647,893
FILING DATE: 30-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 375,731
FILING DATE: 10-JUL-1989
APPLICATION NUMBER: 221,687
FILING DATE: 20-JUL-1988
SEQ ID NO:3
LENGTH: 23
5451527-3

Alignment Scores:
Pred. No.: 320 Length: 23
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 6 Gaps: 0

US-09-745-506-74 (1-1553) x 5451527-3 (1-23)

OY 407 GCACCTCCTGCATCCTCCT 387
DB 1 AlapropProserleupro 7

RESULT 13
US-09-217-306B-24
Sequence 24, Application US/09217306B
Patent No. 6465220
GENERAL INFORMATION:
APPLICANT: Hassan, Helle
APPLICANT: Clausen, Henrik
APPLICANT: Bennett, Eric P.
TITLE OF INVENTION: Glycosylation Using GalNAc-T4 Transferase
FILE REFERENCE: 8850*1
CURRENT APPLICATION NUMBER: US/09/217,306B
CURRENT FILING DATE: 1998-12-21
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Version 3.1
SEQ ID NO 24
LENGTH: 24
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: MISC FEATURE
OTHER INFORMATION: hcg-beta
US-09-217-306B-24

Alignment Scores:
Pred. No.: 318 Length: 24
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 4 Gaps: 0

US-09-745-506-74 (1-1553) x US-09-217-306B-24 (1-24)

OY 407 GCACCTCCTGCATCCTCCT 387
DB 11 AlapropProserleupro 17

RESULT 14
US-08-485-692-13
Sequence 13, Application US/08485692
Patent No. 5739818
GENERAL INFORMATION:
APPLICANT: BOIME, IRVING
TITLE OF INVENTION: MODIFIED PROTEIN AND PEPTIDE
TITLE OF INVENTION: PHARMACEUTICALS
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W.
City: Washington, D.C.
COUNTRY: USA
ZIP: 20006-1812
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,692
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/049,869
FILING DATE: 20-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: MORASHIGE, KATE H.
REGISTRATION NUMBER: 29,959

REFERENCE/DOCKET NUMBER: 29500-20030.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
TELEX: 90-4030
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-485-692-13

Alignment Scores:
Pred. No.: 312 Length: 28
Score: 7.00 Matches: 7
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 1 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-485-692-13 (1-28)

QY 407 GCACCTCCTCATCTCCT 387
DB 6 AlaProProSerLeuPro 12

RESULT 15
US-08-419-519-13
Sequence 13, Application US/08419519
Patent No. 5792460
GENERAL INFORMATION:
APPLICANT: BOIME, IRVING
TITLE OF INVENTION: MODIFIED PROTEIN AND PEPTIDE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W.
CITY: Washington, D.C.
COUNTRY: USA
ZIP: 20006-1812
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentia Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/419,519
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/049,869
FILING DATE: 20-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: MORASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29500-20030.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
TELEX: 90-4030
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-419-519-13

Alignment Scores:
Pred. No.: 312 Length: 28
Score: 7.00 Matches: 7

Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 1.41% Indels: 0
DB: 1 Gaps: 0

US-09-745-506-74 (1-1553) x US-08-419-519-13 (1-28)

QY 407 GCACCTCCTCATCTCCT 387
DB 6 AlaProProSerLeuPro 12

Search completed: August 22, 2003, 14:42:19
Job time : 34.5 secs